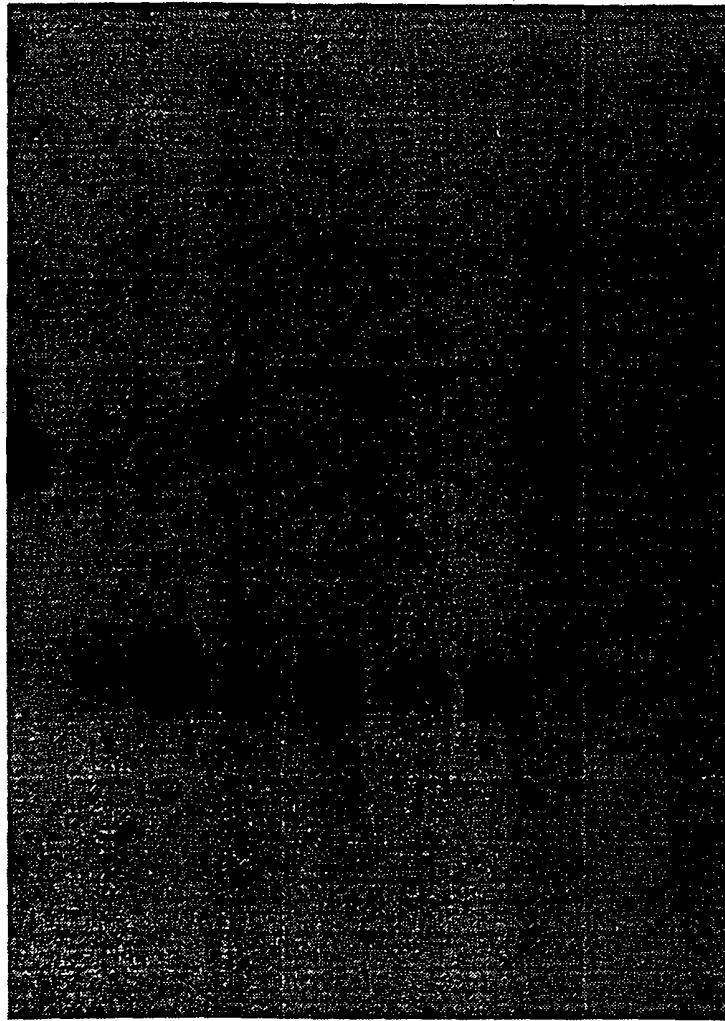


- 28S

- 18S

Fig. 1A



- 28S

- 18S

Fig. 1B

1

Fig. 2

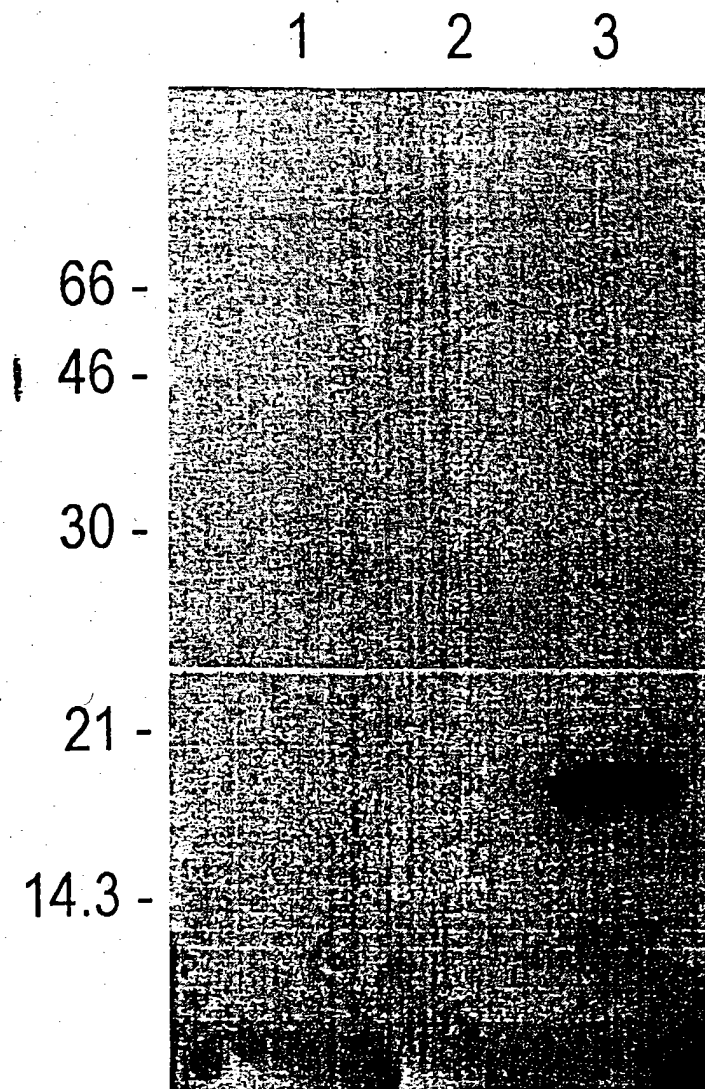


Fig. 3A

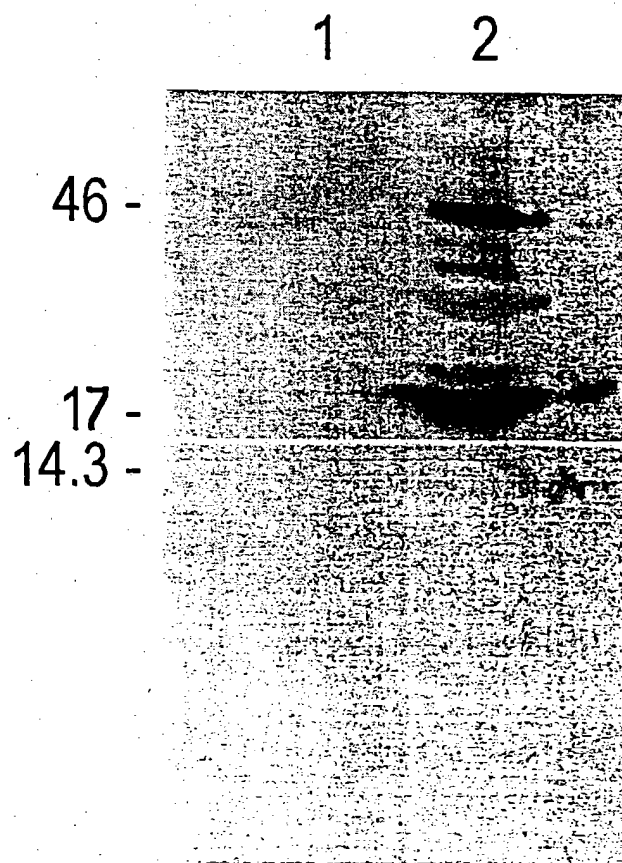


Fig. 3B

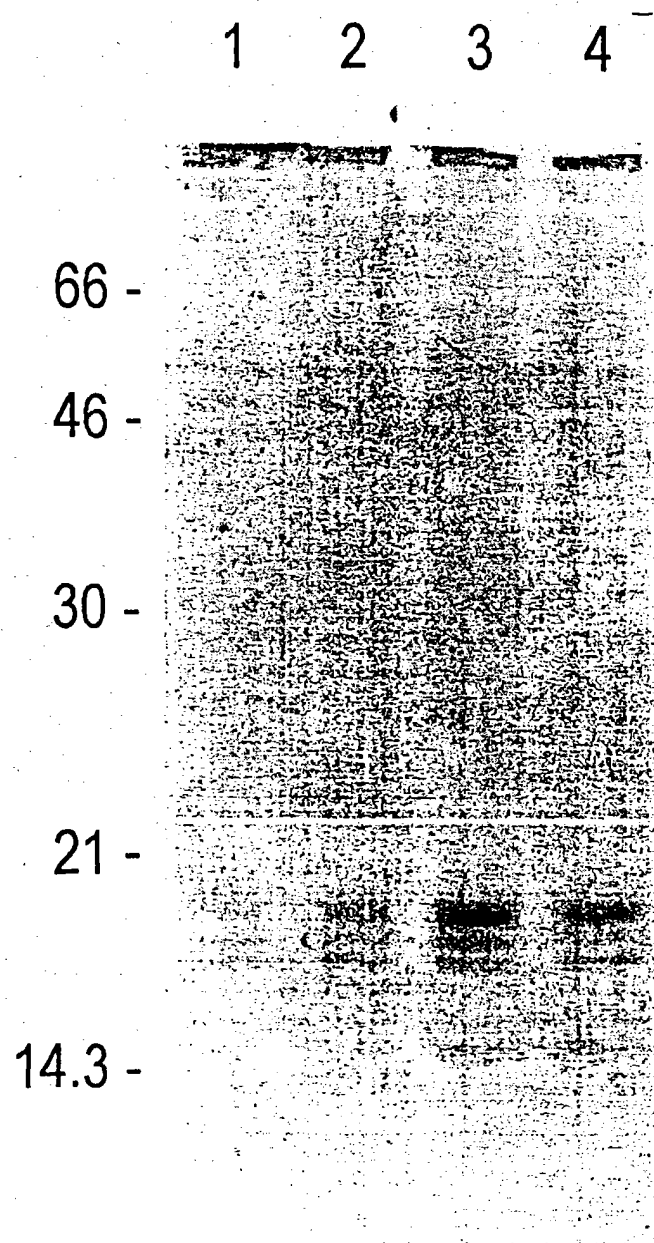


Fig. 3C

GILR	L	K	E	Q	I	R	E	L	L	E	K	N	S	Q	L	E	R	E	N	T	L	L	K	T	L	A
TSC-22	L	K	E	Q	I	K	E	L	I	E	K	N	S	Q	L	E	Q	E	N	D	L	L	K	T	L	A
GCN4	L	E	D	K	V	E	E	L	L	S	K	N	Y	H	L	E	N	E	V	A	R	L	K	K	L	V
CREB	L	E	N	R	V	A	V	L	E	N	Q	N	K	T	L	I	E	E	L	K	A	L	K	D	L	Y
CREM	L	E	N	R	V	A	V	L	E	N	Q	N	K	T	L	I	E	E	L	K	A	L	K	D	L	Y
c-jun	L	E	E	K	V	K	T	L	K	A	Q	N	S	E	L	A	S	T	A	N	M	L	R	E	Q	V

Fig. 4

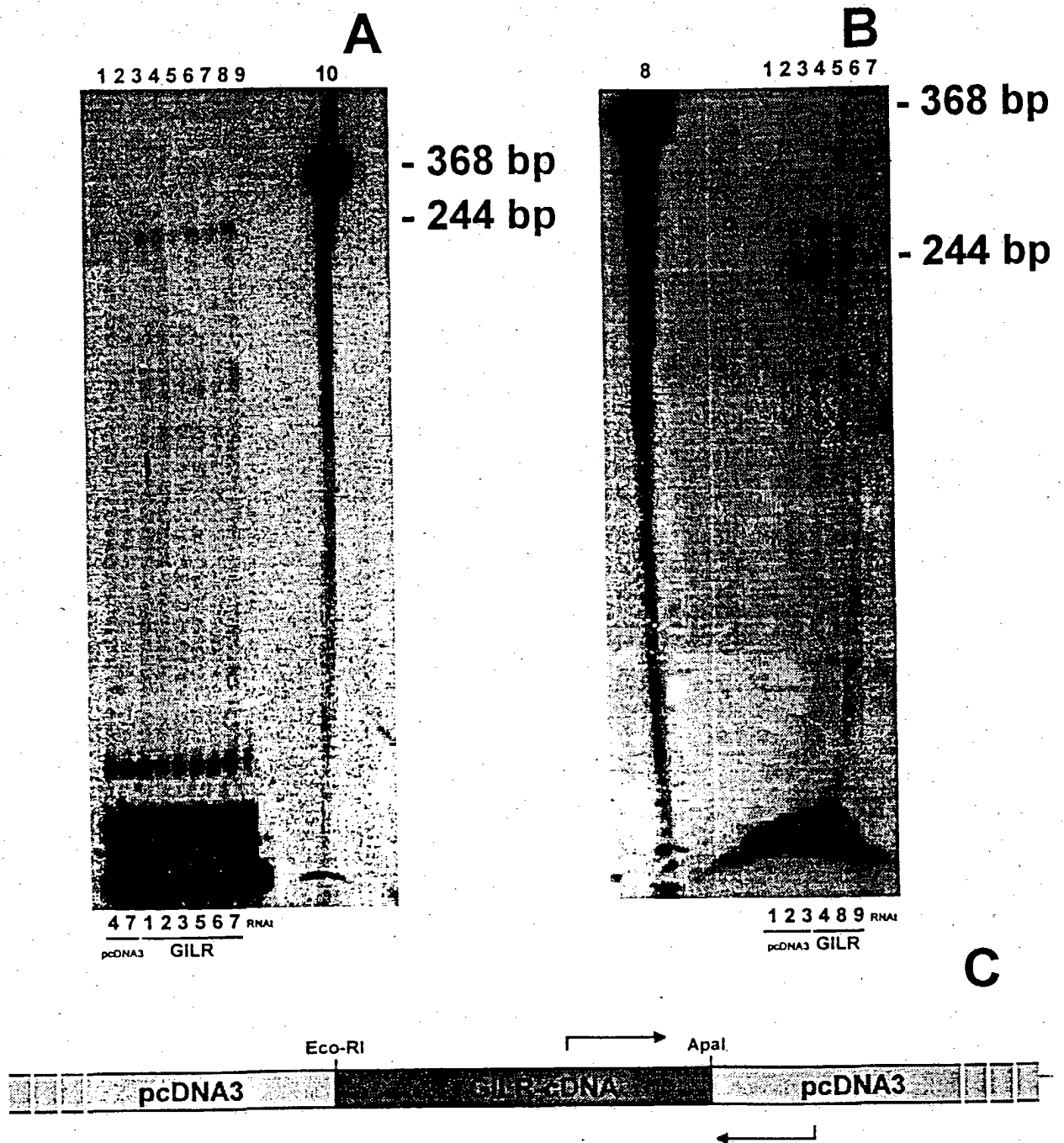


Fig. 5

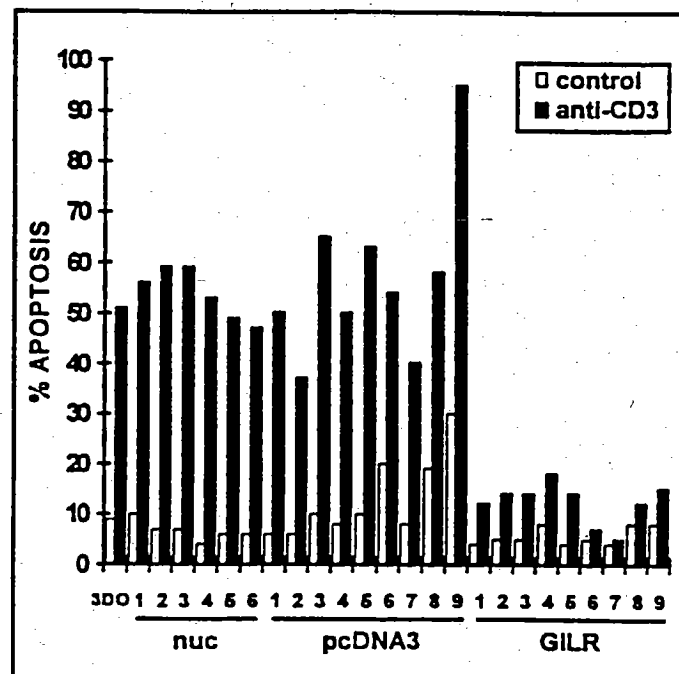


Fig. 6

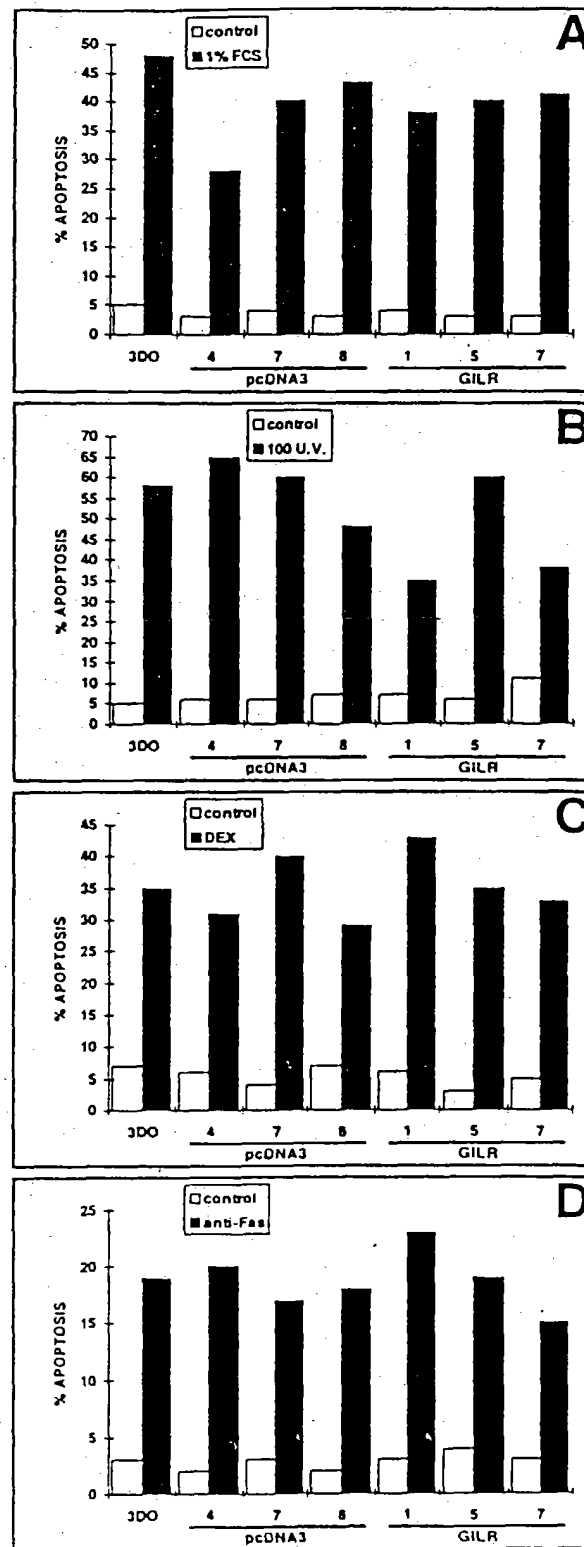


Fig. 7

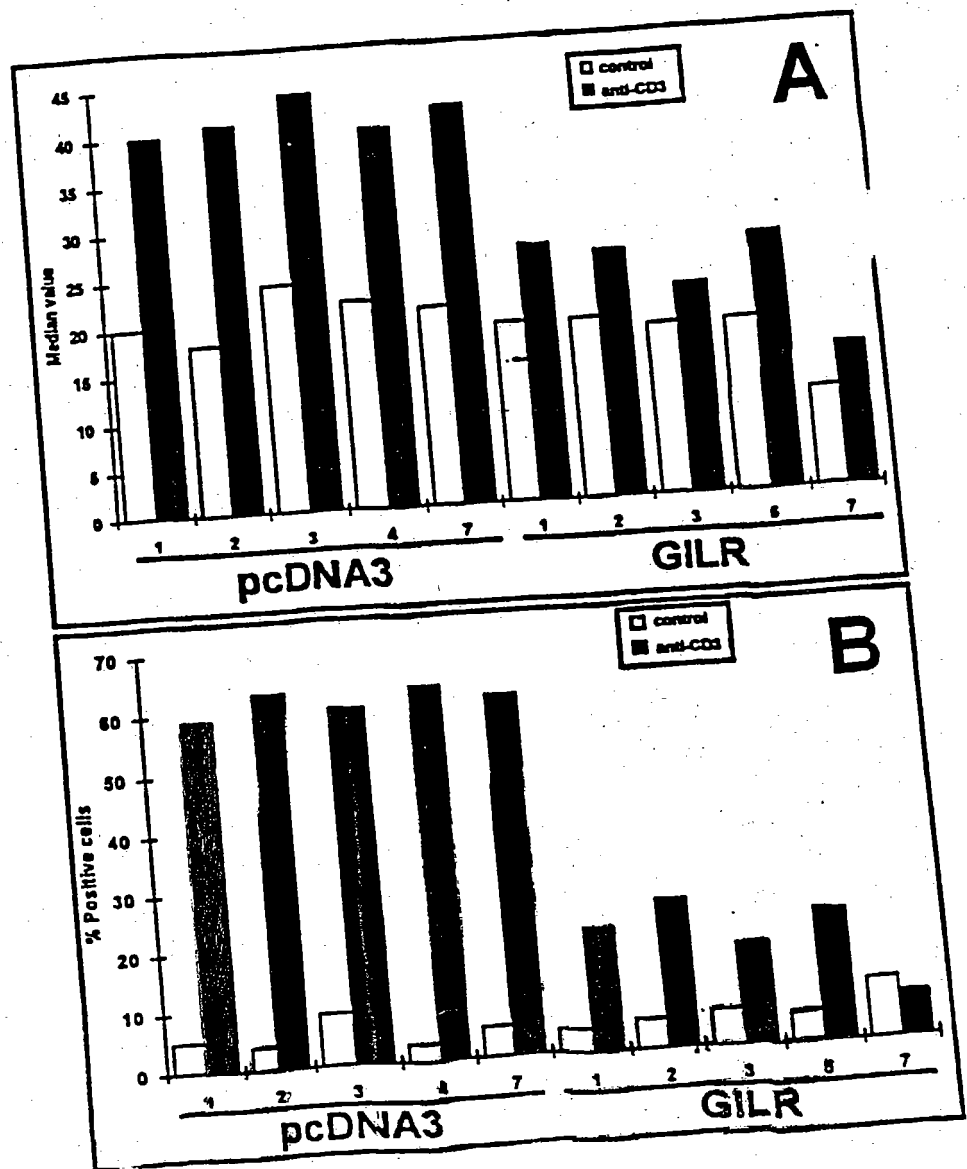


Fig 3

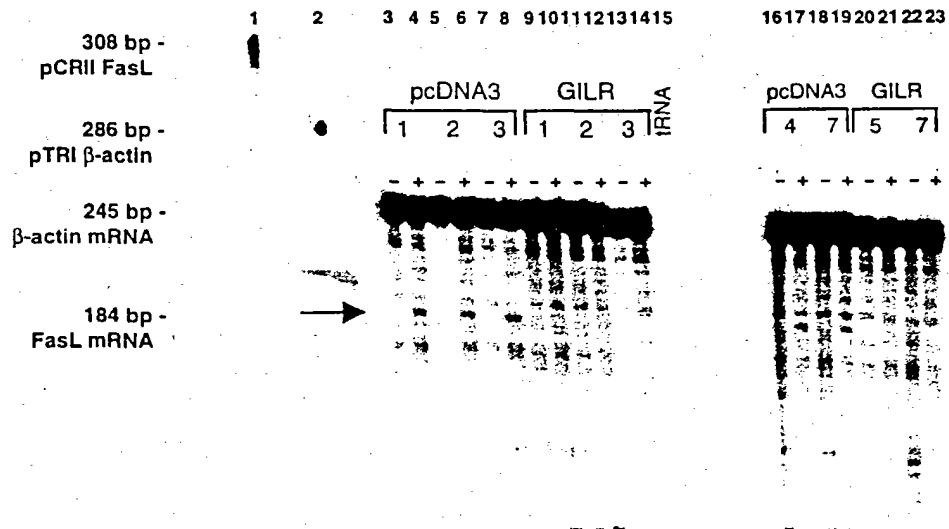
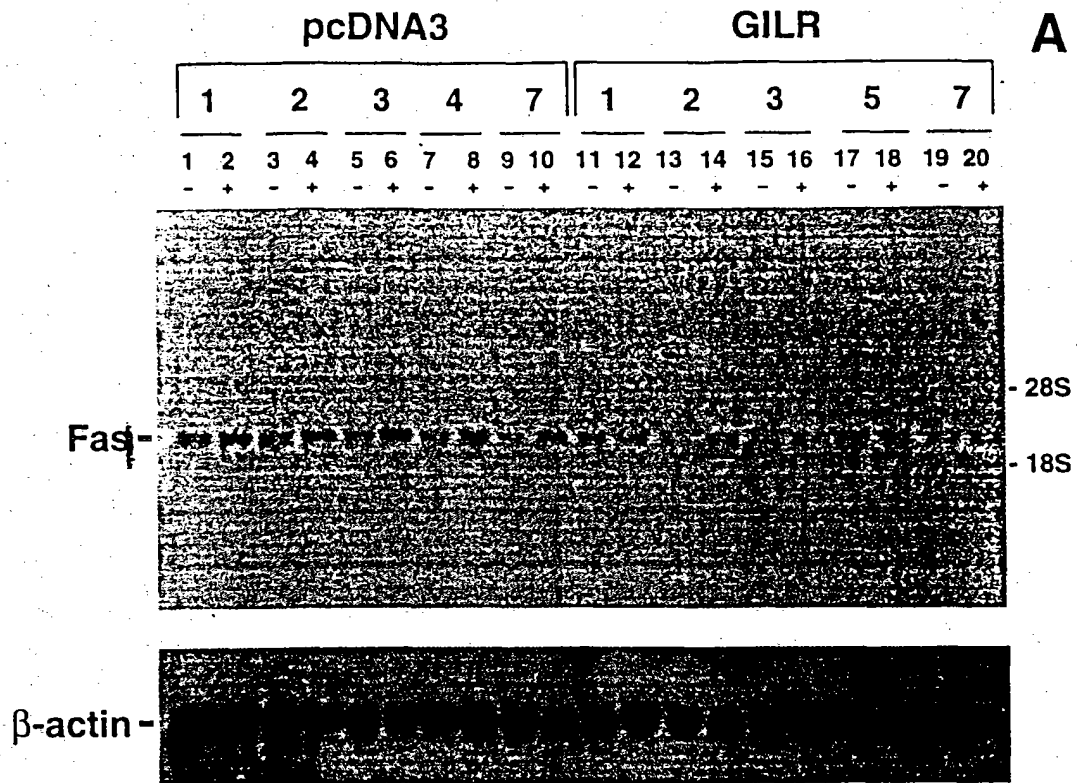
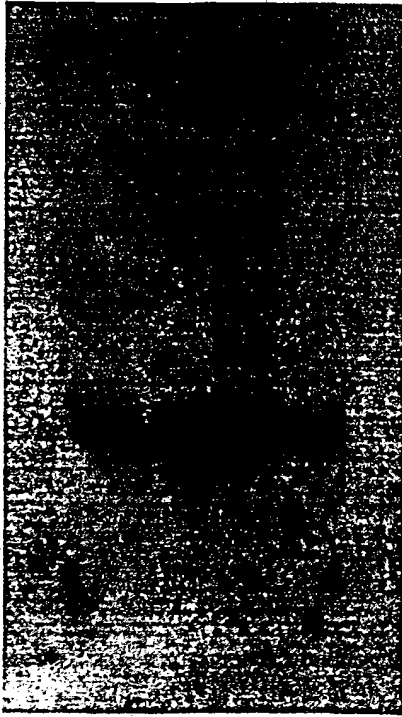
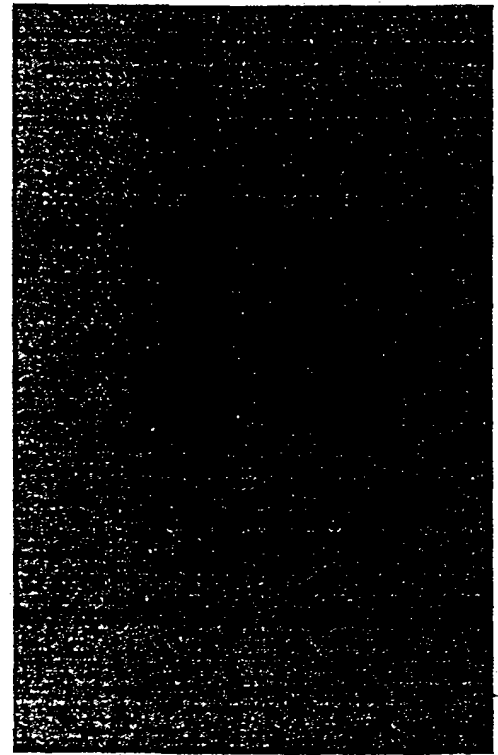


Fig. 9



A



B

Fig. 10

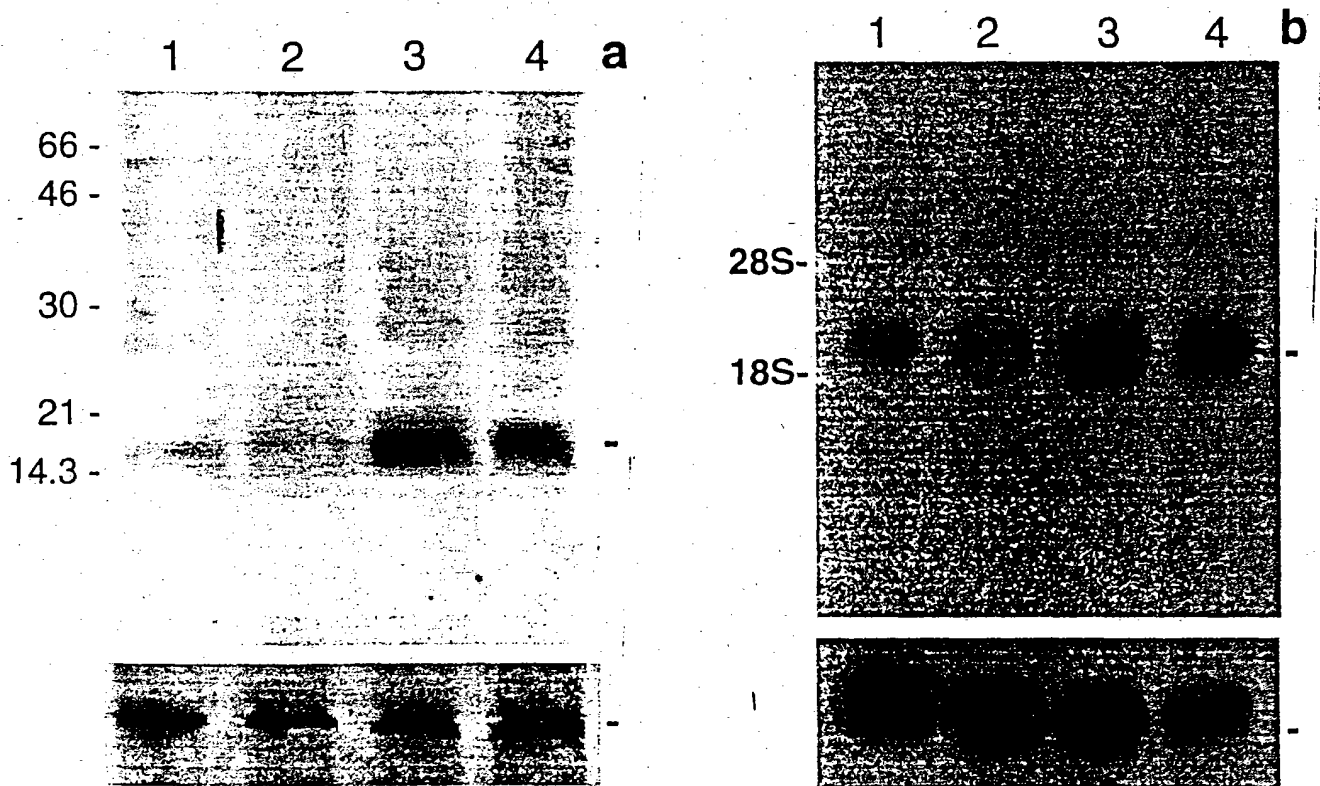


Fig. 11

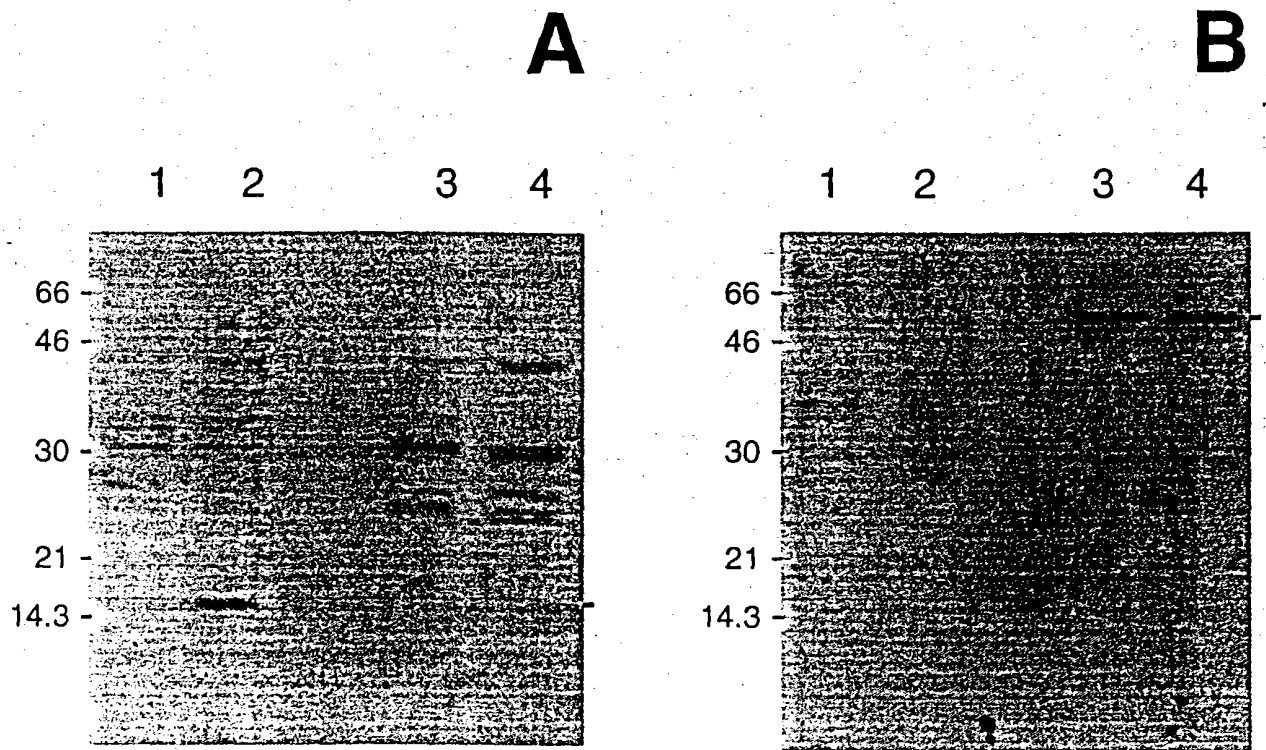


Fig. 12

1 AATTCGGGGGCGGTGGAGTTTGTGACATACGAGGTGACACCCCTCGAGTCACTTCCCTTC
 61 AACTCCAGCTGGAGCGCTGCTTGGCTTTGGGTTCTGCTGACGCTTCGCCCCGCTCCT
 121 AGCCTCAGGGCCGACTCCAGCGCAGAGCCAGCCAGCGCAGCCTGCCAGCAGCCACCC
 181 AGCCGCCCAGCGCCAGCCCGCACGAAACCCGGCCAGAGCTTCTAGCAGCCCGAGCC
 241 ATGAACACCGAAATGTATCAGACCCCATGGAGGTGGCGGTCTACCAGCTGCACAATTC
 MetAsnThrGluMetTyrGlnThrProMetGluValAlaValTyrGlnLeuHisAsnPhe

 301 TCCATCTCCTTCTTCTCTCTGCTTGGAGGGGATGTGGTTTCCGTTAAGCTGGACAAC
 SerIleSerPhePheSerSerLeuLeuGlyGlyAspValValSerValLysLeuAspAsn

 361 ACTGCCTCCGGAGCCAGCGTGGTGGCCATAGACAACAAGATCGAACAGGCCATGGATCTG
 SerAlaSerGlyAlaSerValValAlaIleAspAsnLysIleAspGlnAlaMetAspLeu

 421 GTGAAGAATCATCTGATGTATGCTGTGAGAGAGGAGGTGGAGATCCTGAAGGAGCAGATC
 ValLysAsnHisLeuMetTyrAlaValArgGluGluValGluIleLeuLysGluGlnIle

 481 CGAGAGCTGGTGGAGAAGAACTCCAGCTAGAGCGTGAGAACACCCTGTTGAAGACCCTG
 ArgGluLeuValGluLysAsnSerGlnLeuGluArgGluAsnThrLeuLeuLysThrLeu

 541 GCAA¹CCCAGAGCAGCTGGAGAAGTTCCAGTCCTGTCTGAGCCCTGAAGAGCCAGCTCCC
 AlaSerProGluGlnLeuGluLysPheGlnSerCysLeuSerProGluGluProAlaPro

 601 GAATCCCCACAAGTGCCCGAGGCCCCCTGGTGGTTCTGCGGTGTAAGTGGCTCTGTCTCA
 GluSerProGlnValProGluAlaProGlyGlySerAlaVal *

 661 GGGTGGGCAGAGCCACTAACTTGTFTTACCTAGTTCTTTCCAGTTTGTFTTTGGCTCCC
 721 CAAGCATCATCTCACGAGGAGAACTTTACACCTAGCACAGCTGGTGCCAAGAGATGTCCT
 781 AAGGACATGGCCACCTGGGTCCACTCCAGCGACAGACCCCTGACAAGAGCAGGTCTCTGG
 841 AGGCTGAGTTGCATGGGGCCTAGTAACACCAAGCCAGTGAGCCTCTAATGCTACTGCGCC
 901 CTGGGGGCTCCCAGGGCCTGGGCAACTTAGCTGCAACTGGCAAAGGAGAAGGGTAGTTTG
 961 AGGTGTGACACCAGTTTGCTCCAGAAAGTTTAAGGGGTCTGTTTCTCATCTCCATGGACA
 1021 TCTTCAACAGCTTCACCTGACAACGACTGTTCCCTATGAAGAAGCCACTTGTGTTTTAAGC
 1081 AGAGGCAACCTCTCTCTCTCTGTTTCGTGAAGGCAGGGGACACAGATGGGAGAGAT
 1141 TGAGCCAAGTCAGCCTTCTGTTGGTTAATATGGTATAATGCATGGCTTTGTGCACAGCCC
 1201 AGTGTGGGATTACAGCTTTGGGATGACCGCTTACAAAGTTCTGTTTGGTTAGTATTGGCA
 1261 TAGTTTTTCTATATAGCCATAAATGCGTATATATACCCATAGGGCTAGATCTGTATCTTA
 1321 GTGTAGCGATGTATACATATACACATCCACCTACATGTTGAAGGGCCTAACCAGCCTTG
 1381 GAGTATTGACTGGTCCCTTACCTCTTATGGCTAAGTCTTTGACTGTGTTCAATTACCAAG
 1441 TTGACCCAGTTTGTCTTTTAGGTTAAGTAAGAACTCGAGAGTAAAGGCAAGGAGGGGGC
 1501 CAGCCTCTGAATGCGGCCACGGATGCCTTGCTGCTGCAACCCTTTCCCAGCTGTCCACT
 1561 GAAACGTGAAGTCCTGTTTGAATGCCAAACCCACCATTCACTGGTGCTGACTACATAGA
 1621 ATGGGTTGAGAGAAGATCAGTTTGGGCTTCACAGTGTCATTTGAAAAAGCGTTTTTGT
 1681 TGTTTTGAATTATTGTGGAAAACTTTCAAGTGAACAGAAGGATGGTGTCTACTGTGGAT
 1741 GAGGGATGAACAAGGGGATGGCTTTGATCCAATGGAGCCTGGGAGGTGTGCCCAGAAAGC
 1801 TTGTCTGTAGCGGGTTTTGTGAGAGTGAACACTTTCCACTTTTTGACACCTTATCCTGAT
 1861 GTATGGTTCCAGGATTTGGATTTTGAATTTCCAAATGTAGCTTGAAATTTCAATAAACTT
 1921 TGCTCTGTTTTTCTAAAAATAAAAA

Fig. 13

1

10

```

614 GTGTAAGTGGCTCTGTCCCTTAGGGTGGGCAGAGCCAC..ATCTTGTTCTA 661
|||||
640 gtgtaagtggctctgtcctcaggggtgggcagagccactaaacttgtttta 689
.
662 CCTAGTTCTTTCCAGTTTGTGTTTTGGCTCCCCAAGGGTCATCTCATGTGG 711
|||||
690 cctagttcctttccagtttggttttggtccccaagcatcatctcacgagg 739
.
712 AGAACTTTACACCTAACATAGCTGGTGCCAAGAGATGTCCAAGGACATG 761
|||||
740 agaactttacacctagcacagctggtgccaagagatgtcctaaggacatg 789
.
762 CCCATCTGGGTCCACTCCAGTGACAGACCCCTGACAAAGAGCAGGTCTCT 811
|||
790 gccacctgggtccactccagcgacagaccctgac.aagagcaggtctct 838
.
812 GGAGACTAAGTTGCATGGGGCCTAGTAACACCAAGCCAGTGAGCCTGTCTG 861
||||
839 ggaggctgagttgcatggggcctagtaacaccaagccagtgagcctctaa 888
.
862 TGTCACCGGGCCCTGGGGGCTCCCAGGG.CTGGGCAACTTAGTTACAGCT 910
||
889 tgctactgcgccctgggggctcccagggcctgggcaacttagctgcaact 938
.
911 GACCAAGGAGAAAGTAGTTTTGAGATGTGATGCCAGTGTGCTCCAGAAAG 960
||
939 ggcaaaggagaagggtagtttgaggtgtgacaccagtttgctccagaaag 988
.
961 TGTAAGGGGTCTGTTTTTCATTTCCATGGACATCTTCCACAGCTTCACCT 1010
|
989 tttaaggggtctgtttctcatctccatggacatcttcaacagcttcacct 1038
.
1011 GACAATGACTGTTTCCTATGAAGAAGCCACTTGTGTTCTAAGCAGAAGCAA 1060
||||
1039 gacaacgactgttcctatgaagaagccacttggtgttttaagcagaggcaa 1088
1061 CCTCTCTCTTCTTCTCTGTCTTTTCCAGGCAGGGG.CAGAGATGGGAGA 1109
||||
1089 cctctctcttc.tcctctgtttcgtgaaggcaggggacacagatgggaga 1137
.
1110 GATTGAGCCAAATGAGCCTTCTGTTGGTTAATACTGTATAATGCATGGCT 1159
||||
1138 gattgagccaagtcagccttctggttggttaatatggtataatgcatggct 1187
.
1160 TTGTGCACAGCCCAGTGTGGGGTTACAGCTTTGGGATGACTGCTTATAAA 1209
||||
1188 ttgtgcacagcccagtggtgggattacagctttgggatgaccgcttacaaa 1237
.
1210 GTTCTGTTTGGTTAGTATTGGCATCGTTTTTCTATATAGCCAT.AATGCG 1258
||||
1238 gttctgtttggttagtattggcatagttttctatatatagccataaatgcg 1287
.
1259 TATATATACCCATAGGGCTAGATCTATATCTTAGGGTAGTGATGTATACA 1308
||||
1288 tatatatacccatagggctagatctgtatcttagtgtagcgatgtatata 1337

```

Fig. 14 (cont)

Fig. 14 (Cont.)

Fig. 14 (Cont.)

```

mG  1 MNTEMYQTPMEVAVYQLHNFSTSFSSLLGGDVVSVKLDNSASGASVVAL  50
hg  1 MNTEMYQTPMEVAVYQLHNFSSISFFSSLLGGDVVSVKLDNSASGASVVAI  50
      =====
ht  2 KSQWCRPVAMD LGVYQLRHFSISFLSSLLGTENASVRLDNSSSGASVVAI  51
      =====

mG  51 DNKIEQAMD LVKNHLMYAVREEVEVLKEQIRELLEKNSQLERENTLLKTL 100
hg  51 DNKIEQAMD LVKNHLMYAVREEVEILKEQIRELVEKNSQLERENTLLKTL 100
      =====
ht  52 DNKIEQAMD LVKSHLMYAVREEVEVLKEQIKELIEKNSQLEQENNLLKTL 101
      =====

hd  1          MDLVKNHLMYAVREEVEILKEQIRELVEKNSQLERENTLLKTL  41
      =====

mG 101 ASPEQLEKFQSRLSPEEPAPEAPETPETPEAPGGS AV* 138
hg 101 ASPEQLEKFQSCLSPEEPAPES...PQVPEAPGGS AV* 135
      =====
ht 102 ASPEQLAQFQAQLQTGSPATTQPQGTTPPAQPASQSGGPTA* 145
      =====
hd  42 ASPEQLEKFQSCLSPEEPAPES...PQVPEAPGGS AV*
      =====

```

Fig. 15

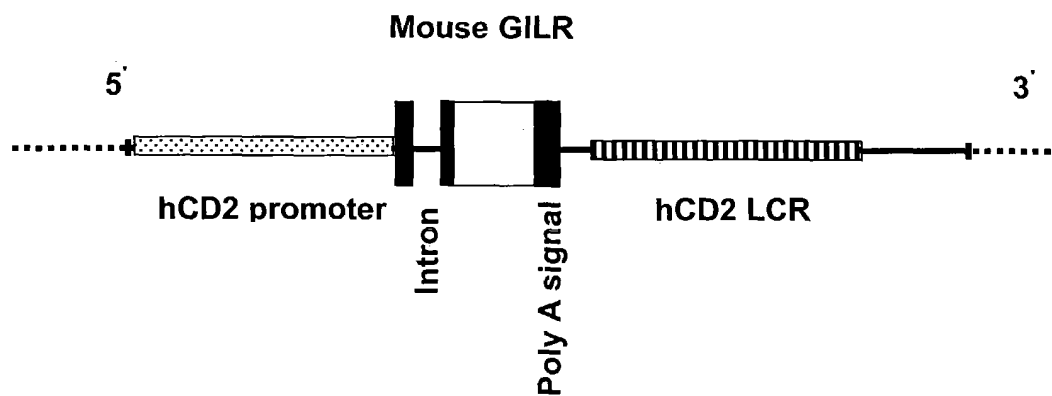


FIG. 16A

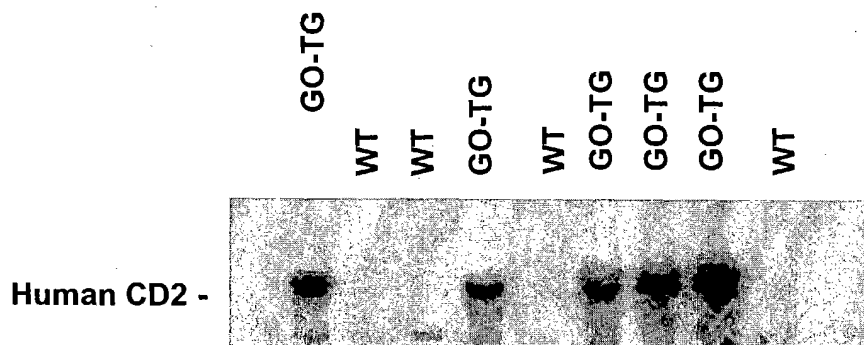


FIG. 16B

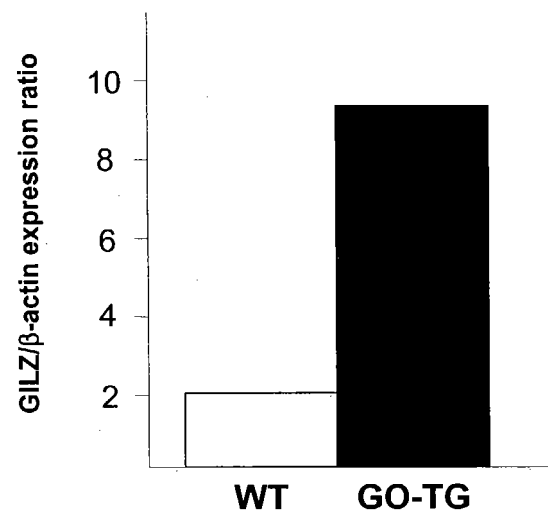


FIG. 17A

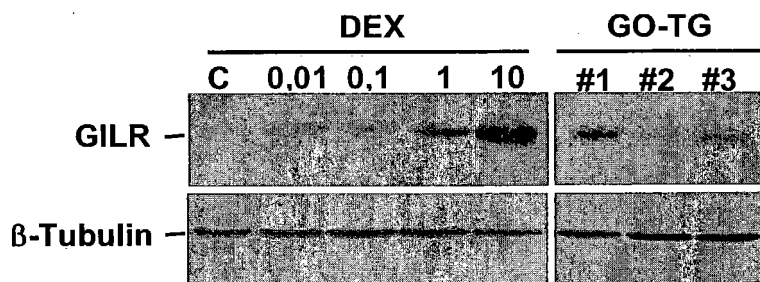


FIG. 17B

FIG. 18A

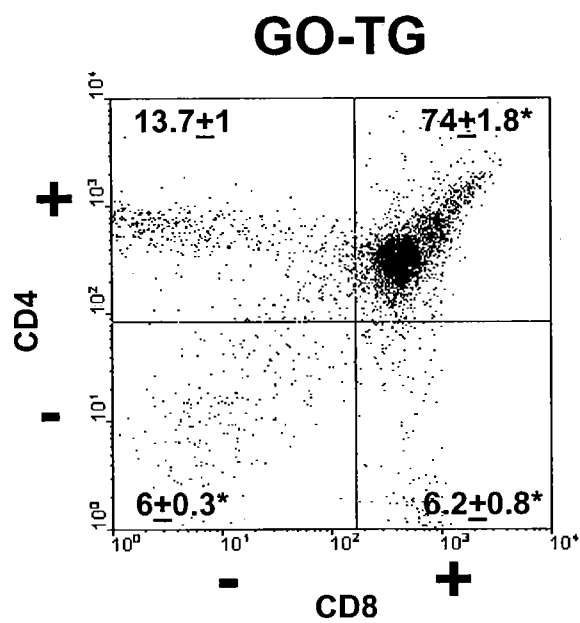
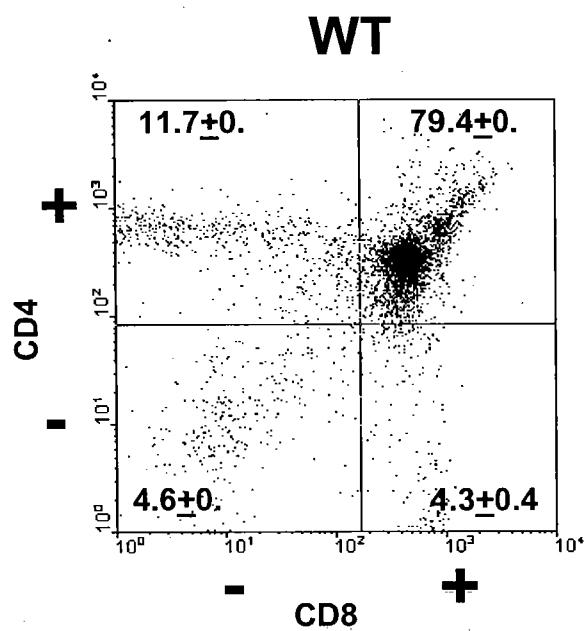


FIG. 18B

WT

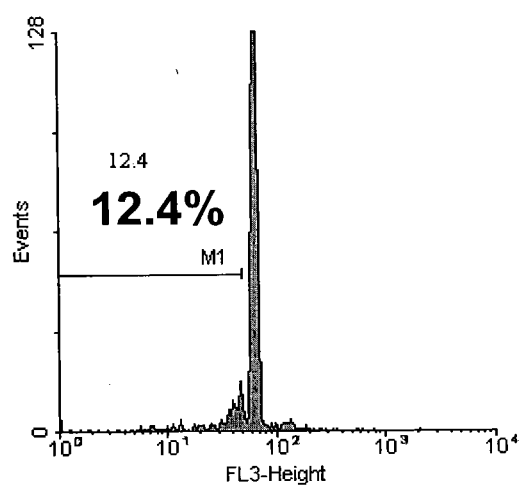


FIG. 19A

GO-TG

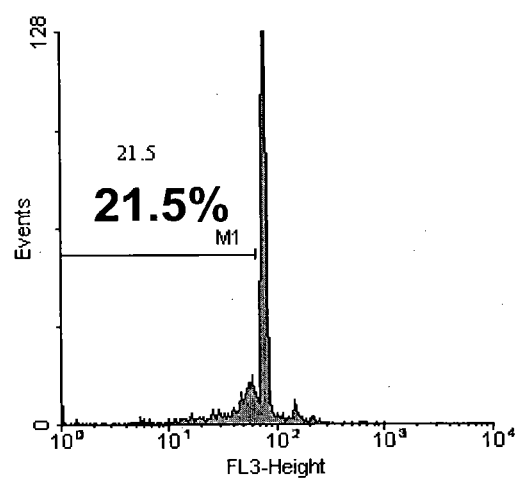


FIG. 19B

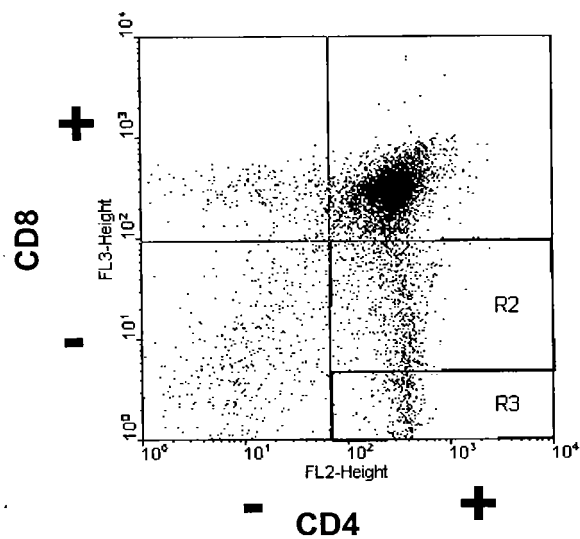


FIG. 20A

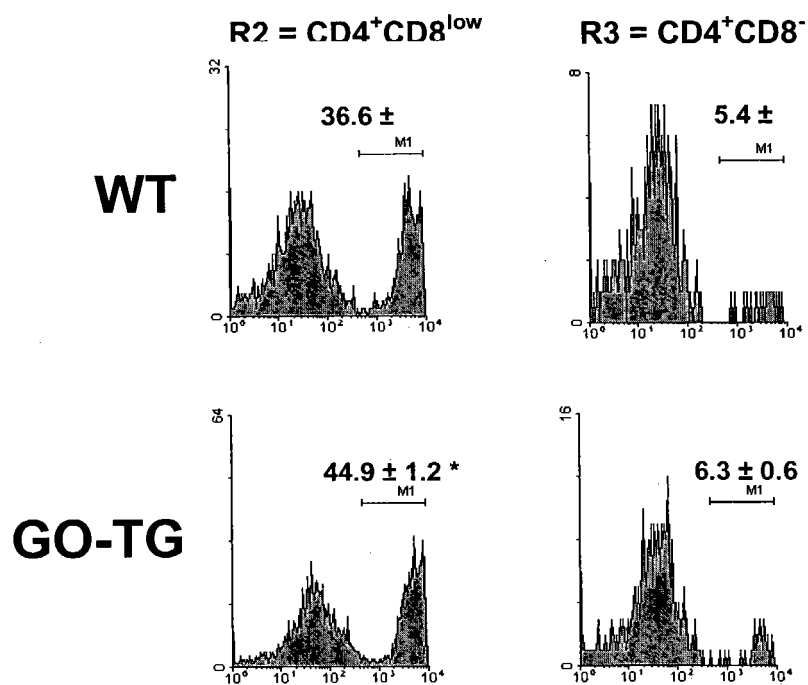


FIG. 20B

FIG. 21A

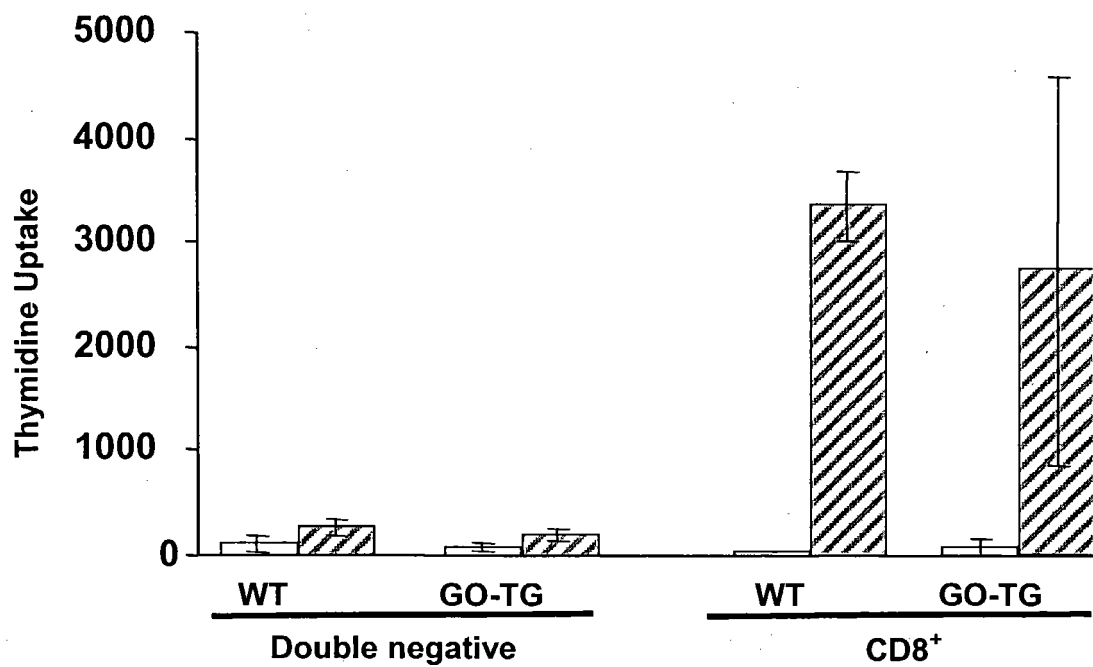
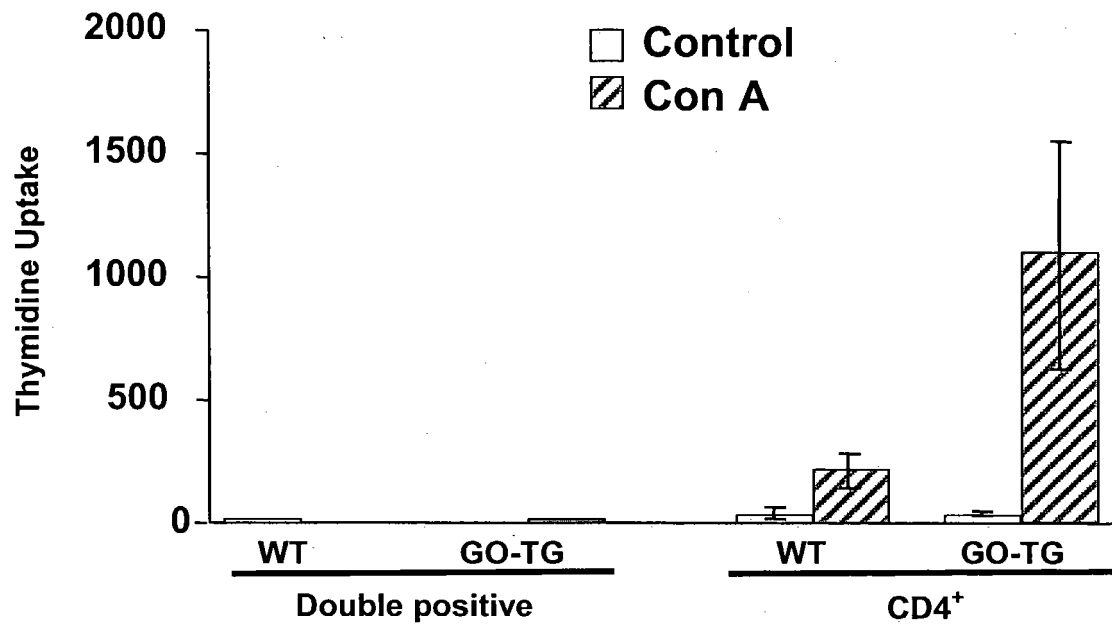


FIG. 21B

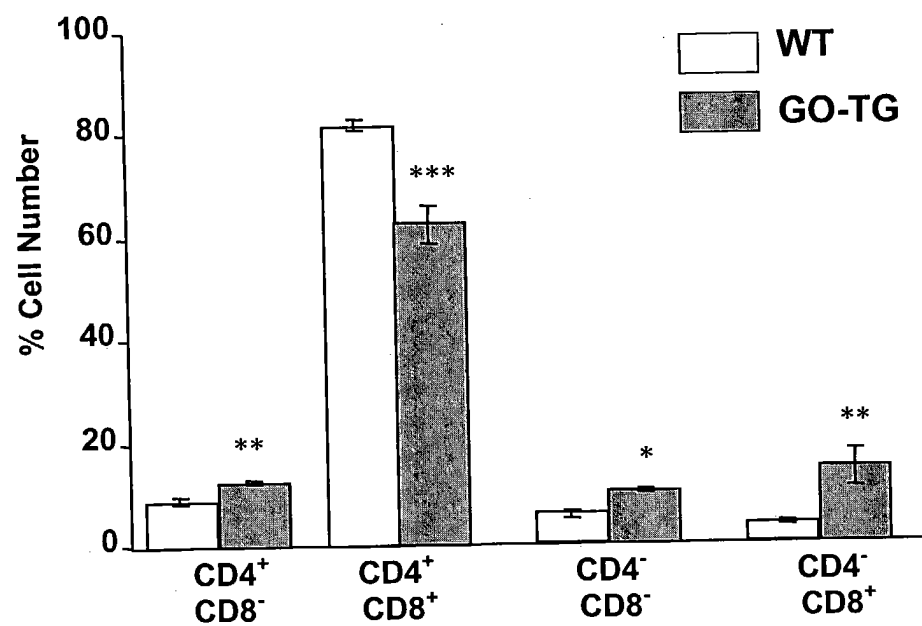


FIG. 22

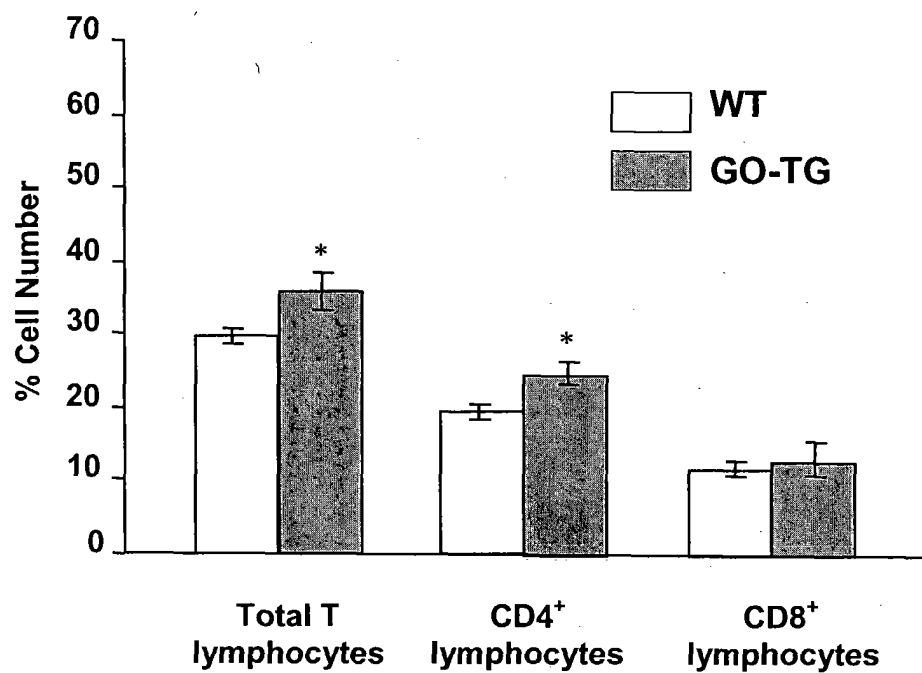


FIG. 23

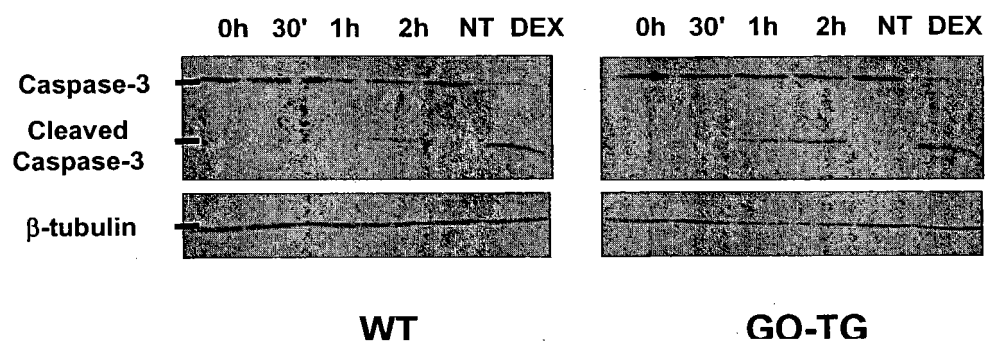


FIG. 24